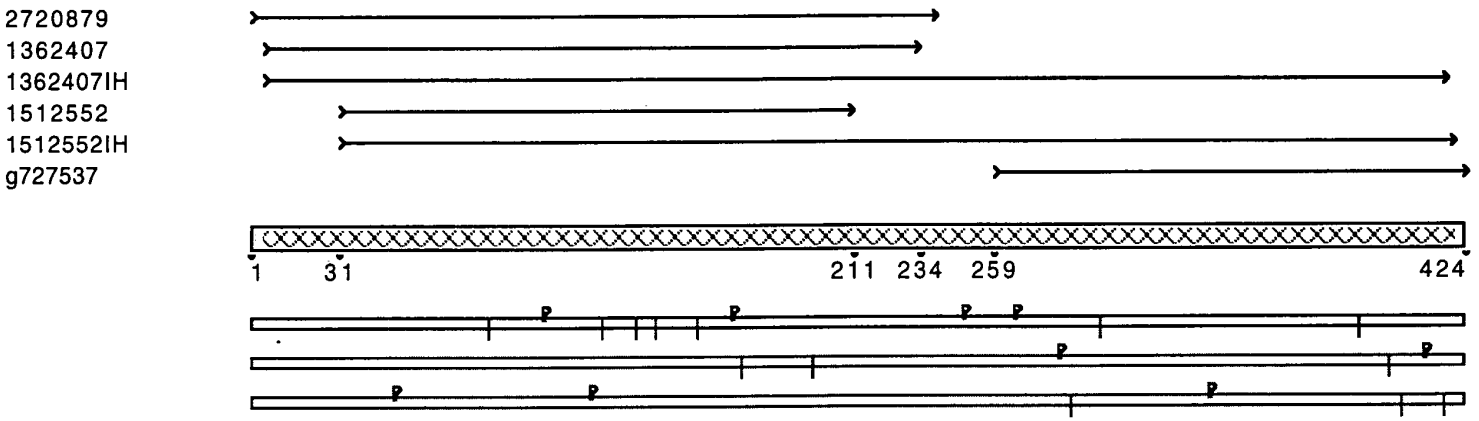
Figure 1-A

```
>2720879        GGCCACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT
>1362407            ACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT
>1362407IH          ACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT
>1512552                                          CAGGAGCGC AGTGGCCACT
>1512552IH                                       CAGGAGCGC AGTGGCCACT
Consensus       GGCCACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT

>2720879        ATGGGGTCTG GGCTGCCCCT TGTCCTCCTC TTGACCCTCC TTGGCAGCTC
>1362407        ATGGGGTCTG GGCTGCCCCT TGTCCTCCTC TTGACCCTCC TTGGCAGCTC
>1362407IH      ATGGGGTCTG GGCTGCCCCT TGTCCTCCTC TTGACCCTCC TTGGCAGCTC
>1512552        ATGGGGTCTG GGCTGCCCCT TGTCCTCCTC TTGACCCTCC TTGGCAGCTC
>1512552IH      ATGGGGTCTG GGCTGCCCCT TGTCCTCCTC TTGACCCTCC TTGGCAGCTC
Consensus       ATGGGGTCTG GGCTGCCCCT TGTCCTCCTC TTGACCCTCC TTGGCAGCTC

>2720879        ACATGGAACA GGGCCGGGTA TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1362407        ACATGGAACA GGGCCGGGTA TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1362407IH      ACATGGAACA GGGCCGGGTA TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1512552        ACATGGAACA GGGCCGGGTA TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1512552IH      ACATGGAACA GGGCCGGGTA TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
Consensus       ACATGGAACA GGGCCGGGTA TGACTTTGCA ACTGAAGCTG AAGGAGTCTT

>2720879        TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAN
>1362407        TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
>1362407IH      TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
>1512552        TTCTGACAAG TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
>1512552IH      TTCTGACAAG TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
Consensus       TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG

>2720879        :TCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA
>1362407        :TCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCA
>1362407IH      CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA
>1512552        CTCTGCCTCC T
>1512552IH      CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA
Consensus       CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA

>1362407IH      TGCAAGATCT CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC
>1512552IH      TGCAAGATCT CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC
>g727537                T CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC
Consensus       TGCAAGATCT CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC

>1362407IH      CTGTGTCCTT CTTGGCCCGG GCTTTTGGGC CGGGGATGCA GGAGGCAGGC
>1512552IH      CTGTGTCCTT CTTGGCCCGG GCTTTTGGGC CGGGGATGCA GGAGGCAGGC
>g727537        CTGTGTCCTT CTTGGCCCGG GCTTTTNGGC GGGGAATGCA GGAGGCAGGC
Consensus       CTGTGTCCTT CTTGGCCCGG GCTTTTGGGC CGGGGATGCA GGAGGCAGGC

>1362407IH      CCCGACCCTG TCTTTCAGCA GGCCCCCACC CTCCTGAGTG GCAATAAATA
>1512552IH      CCCGACCCTG TCTTTCAGCA GGCCCCCACC CTCCTGAGTG GCAATAAATA
>g727537        CCCGACCCTN TCTTTCAGCA GGCCCCCACC CTNCTGAGTN GCAATAAATA
Consensus       CCCGACCCTG TCTTTCAGCA GGCCCCCACC CTCCTGAGTG GCAATAAATA
```

Figure 1-B

```
>1362407IH      AAATTCGGTA TGCTTGA
>1512552IH      AAATTCGGTA TGCTGAATTC
>g727537        AAATTCGGTA TGCTGAATTC AATA
Consensus       AAATTCGGTA TGCTGAATTC AATA
```

Figure 2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1362407IH
1512552
1512552IH
g727537
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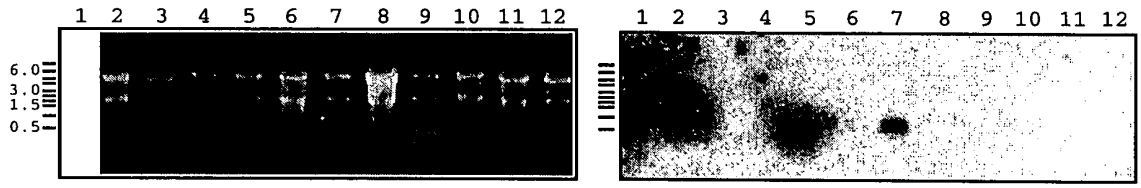
31
211
234
259
424
P
P
P
P
P
P
P
P
P

Figure 3



Lane	Tissue	Lane	Tissue
1	Bladder	7	Lung
2	Brain	8	Ovary
3	Breast	9	Placenta
4	Colon	10	Prostate
5	Kidney	11	Spleen
6	Liver	12	Testis